

Figure 1. Diagram of the paralogue cluster

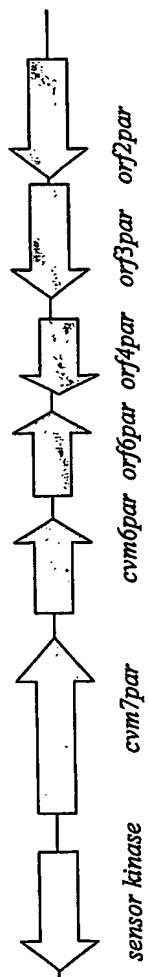


Fig.2 Orientation of *cvm7* to published *cvm* cluster

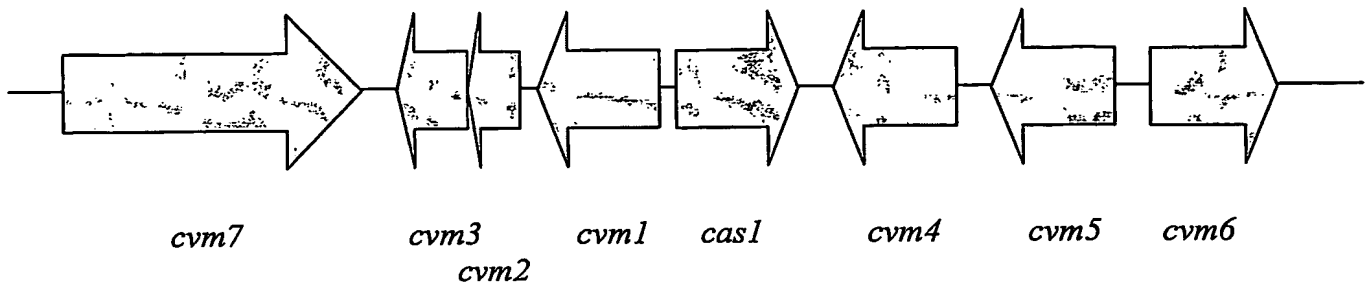


Fig 3. Annotated sequence of the paralogue cluster

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      | 10 | 20 | 30 | 40 | 50 | 60 | 70 | 80
1  ccatgggagc agcatgcag tgcgcctccc cgccgccat gcgctagct ggtagtcccc ctgcgggtg ccgaccgccg 80
      | 81 | 90 | 100 | 110 | 120 | 130 | 140 | 150 | 160
81  gggcggtccc ggtgcgccg gccggatcta gtcgggtgtgc tccgacggtg cctgctgggt gaggggcagt gtcaggcgga 160
      | 161 | 170 | 180 | 190 | 200 | 210 | 220 | 230 | 240
161  tggtagttcc cgcgcggggc gggctgtgca gccgcagttg gccgcagagt gcctccacc ggctcgtgag gccgacgagg 240
      | 241 | 250 | 260 | 270 | 280 | 290 | 300 | 310 | 320
241  cccgagcccc ggcagggggc ggcgccaccg cggccgctgt cgcggatgcc gacgtggagc cgtccgtccc gggtagccac 320
      | 321 | 330 | 340 | 350 | 360 | 370 | 380 | 390 | 400
321  atggacgtcg acgacggttg caccggagtg cttggcggcg ttggtcaggg cctcggagac ggcgtagtac gcggcggtct 400
      | 401 | 410 | 420 | 430 | 440 | 450 | 460 | 470 | 480
401  cgaccggttc ggggtggcgt tccccgtct ggatgtcgag ccggaccggg atggcggagc gccgggccag ggccttgagc 480
      | 481 | 490 | 500 | 510 | 520 | 530 | 540 | 550 | 560
481  gccgggggga gtccgcctc ggcgagtagc gccgggtgga tgccccgggc gacctcccg agttcgtcga cggcggcggc 560
      | 561 | 570 | 580 | 590 | 600 | 610 | 620 | 630 | 640
561  cagcccgctg gtcacctcgt cgaagtgcg gatcagctcg tcggcgctga gcggcacca cagttgcacg gtgcgcacc 640
      | 641 | 650 | 660 | 670 | 680 | 690 | 700 | 710 | 720
641  gcagcgccag ggagaccag cgtctgttgg ggcgtcgtg caggtcgcgt tcgatacggc ggcgggcggt gtcggcggcg 720
      | 721 | 730 | 740 | 750 | 760 | 770 | 780 | 790 | 800
721  gcgacgaccc gggcccgtag cgcggtgagg gccgcctgcg tctccgcgtt ggcgatggcg gtggccacca gttcgggtga 800
      | 801 | 810 | 820 | 830 | 840 | 850 | 860 | 870 | 880
801  gccggccagc cgttcctcgg tgtccgacgg catcggttg tcgttcacg acgcacgct gagcgcgcc cagagttgtc 880
      | 881 | 890 | 900 | 910 | 920 | 930 | 940 | 950 | 960
881  cgtcgacgtt gatcggcatg cacaccgttg cgcggaatcc ccaactcttg ccgacgacgg aggcggggcc cgaggacacg 960
      | 961 | 970 | 980 | 990 | 1000 | 1010 | 1020 | 1030 | 1040
961  gccgcgtagt cgtcgatccg cgcggggcag cccgactcga acaccaggtt gtgcacattc cggcgcggcg gcggtacctg 1040
      | 1041 | 1050 | 1060 | 1070 | 1080 | 1090 | 1100 | 1110 | 1120
1041  gataccggcg ggaataatcac gccgggtcct ggtccaggcg gcgacataca ggcgggttcc gttgggctcg taacggccga 1120
      | 1121 | 1130 | 1140 | 1150 | 1160 | 1170 | 1180 | 1190 | 1200
1121  ggacgcgaaa gtcggccgag agagctgtc cggcctcggc ggcgaccgcg gcgaacacct ccttcggcgg tccgccccgc 1200
      | 1201 | 1210 | 1220 | 1230 | 1240 | 1250 | 1260 | 1270 | 1280
1201  gcgaccaggg tcgccacgcg ccgcagcgcc gcctgtctct cggcgggccc ccgcagctcc acacgtgcct ggggtgtcgc 1280

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36

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3e

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12881 CGTCCACGA TCCGAGGAGA ATCCATGGAT GTCTGGCGG CGTTGGAGCG CAAGCCGAG CTGAATCTTT TCCCATCGA 12960
12961 GAACCGGCTG TCGCCGGCG CCAAGTGCCG GCTGGCCACC GACGCCGTCA ACCGCTATCC GTACTCCAG ACCCGGTGG 13040
13041 CGCTCTACG CGATGTCAG GGGCTGGCG AGGTACGC GTACTGCGAG GACCTGGCCA AGCGTTCTT CCGGGCGCG 13120
13121 CAGCCGGTG TGCAGTCTCT GTCCGGTCTG CACACCATG ACACGTGCT GACCGCCTG ACCCGCGCG GCGGGCGCGT 13200
13201 CCTGGTCTC GCGCGGAGG ACGGGGCCA CTACGCCAG GTGACGATCT GCGGGGCTT CGGCTACGAG CTCGACTTCT 13280
13281 TACCTTCGAC CGCCGGACAC CTGGAGATCG ACT 13313
| 10 | 20 | 30 | 40 | 50 | 60 | 70 | 80